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Application : 09/466	568 Examiner: 1	nchelvey	GAU:	1636
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To: K. Pinkney USPTO 703-308-6642

From: Melissa S. Rones

Comments:

Re: U.S. Application No. 09/466568

Attorney Docket No.: APBI-P16-316

As per your request, enclosed please find a copy of the sequence listing filed on August 12, 2002 in the above-referenced case.

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Date: May 13, 2005

File Symbol: APBI-P16-316

Submitted By: Melissa S. Rones

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Approved for use through 07/31/2003. OMB 0651-0031 U.S. Patent and Trademark Office; U.S. DEPARTMENT OF COMMERCE Under the Paperwork Reduction Act of 1995, no persons are required to respond to a collection of information unless it displays a valid OMB control number. Application Number 09/466568 Filing Date **TRANSMITTAL** December 17, 1999 First Named Inventor **FORM** Gerald R. Crabtree Art Unit 1636 (to be used for all correspondence after initial filing) **Examiner Name** T. A. McKelvey Attorney Docket Number APBI-P16-316 Total Number of Pages in This Submission ENCLOSURES (Check all that apply) After Allowance Communication Fee Transmittal Form Drawing(s) to TC Appeal Communication to Board of Fee Attached Licensing-related Papers Appeals and Interferences Appeal Communication to TC Amendment/Reply Petition (Appeal Notice, Brief, Reply Brief) Petition to Convert to a After Final Proprietary Information Provisional Application Power of Attorney, Revocation Status Letter Affidavits/declaration(s) Change of Correspondence Address X Other Enclosure(s) (please Identify below): **Extension of Time Request** Terminal Disclaimer **Express Abandonment Request** Request for Refund Information Disclosure Statement CD, Number of CD(s) Certified Copy of Priority Landscape Table on CD Document(s) Reply to Missing Parts/ Copy of sequence listing (33 pages) Remarks Incomplete Application Reply to Missing Parts under 37 CFR 1.52 or 1.53 SIGNATURE OF APPLICANT, ATTORNEY, OR AGENT Firm Name ROPES & GRAY LLP Signature Printed name Melissa S. Rones, Ph.D. Reg. No. Date 54,408 May 13, 2005

I hereby certify that this correspondence	is being facsimile transmitted to the Petent and	Trademark Office, facsimile no., on the date
shown below.	1	
Dated:	Signature:	(Ginny Blundell)

SEQUENCE LISTING

(1) GENERAL INFORMATION:

- (i) APPLICANT: Crabtree, Gerald R.
 Schreiber, Stuart L.
 Spencer, David M.
 Wandless, Thomas J.
 Belshaw, Peter
- (ii) TITLE OF INVENTION: REGULATED TRANSCRIPTION OF TARGETED GENES AND OTHER BIOLOGICAL EVENTS
- (iii) NUMBER OF SEQUENCES: 81
- (iv) CORRESPONDENCE ADDRESS:
 - (A) ADDRESSEE: ARIAD Pharmaceuticals, Inc.
 - (B) STREET: 26 Landsdowne Street
 - (C) CITY: Cambridge
 - (D) STATE: Massachusetts
 - (E) COUNTRY: USA
 - (F) ZIP: 02139
 - (v) COMPUTER READABLE FORM:
 - (A) MEDIUM TYPE: Floppy disk
 - (B) COMPUTER: IBM PC compatible
 - (C) OPERATING SYSTEM: PC/DOS/MS/DOS
 - (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
- (vi) CURRENT APPLICATION DATA:
 - (A) APPLICATION NUMBER: US 08/478,386
 - (B) FILING DATE: 07/JUN/1995
 - (C) CLASSIFICATION: 435
- (viii) ATTORNEY/AGENT INFORMATION:
 - (A) NAME: Figg, E. Anthony
 - (B) REGISTRATION NUMBER: 27,195
 - (C) REFERENCE/DOCKET NUMBER: 2054-114A
 - (ix) TELECOMMUNICATION INFORMATION:
 - (A) TELEPHONE: (202) 783-6040
 - (B) TELEFAX: (202) 783-6031

(2)	INFORMATION FOR SEQ ID NO:1:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 14 amino acids (B) TYPE: amino acid (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: peptide	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:	
Me 1	Gly Ser Ser Lys Pro Lys Asp Pro Ser Gln Arg 5 10	
(2	INFORMATION FOR SEQ ID NO:2:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:2:	
G1	AAGTTAA C	13
(2	INFORMATION FOR SEQ ID NO:3:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 11 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
	(ii) MOLECULE TYPE: cDNA	
	(xi) SEQUENCE DESCRIPTION: SEQ ID NO:3:	
T	ACTCAGCG C	1
(INFORMATION FOR SEQ ID NO:4:	
	 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 33 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	

(ii) MOLECULE TYPE: cDNA

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: 6..11
- (D) OTHER INFORMATION: /note= "Sac II restriction site."
- (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: $12..\overline{16}$
 - (D) OTHER INFORMATION: /note= "Kozak sequence."
- (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 17..31
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $17..\overline{33}$
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:4:

CGACACCGCG GCCACC ATG GCC ACA ATT GGA GC Met Ala Thr Ile Gly

- (2) INFORMATION FOR SEQ ID NO:5:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:5:

Met Ala Thr Ile Gly 1 5

- (2) INFORMATION FOR SEQ ID NO:6:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: $12..\overline{27}$
- (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:6:

CGACACTCGA GAGCCCATGA CTTCTGG

27

- (2) INFORMATION FOR SEQ ID NO:7:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4 amino acids

 - (B) TYPE: amino acid(C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..4
 - (D) OTHER INFORMATION: /note= "Translation product of complement of SEQ ID NO:6, bases 9 to 20."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:7:

Ser Trp Ala Leu 1

- (2) INFORMATION FOR SEQ ID NO:8:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 41 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $12..\overline{41}$
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."

(ix) FEATURE:

(A) NAME/KEY: CDS

(B) LOCATION: 9..41

(ix) FEATURE:

(A) NAME/KEY: misc_feature

(B) LOCATION: 28

- (D) OTHER INFORMATION: /note= "A to G."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:8:

CGACACTC GAG CTC TGC TAC TTG CTA GGT GGA ATC CTC TTC Glu Leu Cys Tyr Leu Leu Gly Gly Ile Leu Phe 41

- (2) INFORMATION FOR SEQ ID NO:9:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:9:

Glu Leu Cys Tyr Leu Leu Gly Gly Ile Leu Phe 5 1

- (2) INFORMATION FOR SEQ ID NO:10:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 24 base pairs (B) TYPE: nucleic acid

 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 3..8
 - (D) OTHER INFORMATION: /note= "Eco RI restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 9..24
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 24
 - (D) OTHER INFORMATION: /note= "G to C."

(ix) FEATURE:

(A) NAME/KEY: misc signal

(B) LOCATION: complement (9..11)

(D) OTHER INFORMATION: /note= "Translational stop encoded in complementary strand."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:10:

GCGAATTCTT AGCGAGGGGC CAGC

24

- (2) INFORMATION FOR SEQ ID NO:11:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..4
 - (D) OTHER INFORMATION: /note= "Translational product of complement to SEQ ID NO:10, bases 12 to 23."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:11:

Leu Ala Pro Arg

- (2) INFORMATION FOR SEQ ID NO:12:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 3..8
 - (D) OTHER INFORMATION: /note= "Eco RI restriction."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $12..\overline{17}$
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: complement (9..11)

33

(D) OTHER INFORMATION: /note= "Translational stop signal encoded on complementary strand."

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: 18..33
- (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:12:

GCGAATTCTT AGTCGACGCG AGGGGCCAGG GTC

(2) INFORMATION FOR SEQ ID NO:13:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 4 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: peptide
- (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..4
 - (D) OTHER INFORMATION: /note= "Translational product of complement to SEQ ID NO:12, bases 18 to 29."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:13:

Leu Ala Pro Arg 1

- (2) INFORMATION FOR SEQ ID NO:14:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 25 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 4..9
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 13
 - (D) OTHER INFORMATION: /note= "T to G."

25

(ix) FEATURE:

(A) NAME/KEY: misc_feature

- (B) LOCATION: 4..25
 (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 10..24
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:14:

GGGCTCGAG CTC GGC TAC TTG CTA G Leu Gly Tyr Leu Leu

(2) INFORMATION FOR SEQ ID NO:15:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:15:

Leu Gly Tyr Leu Leu

- (2) INFORMATION FOR SEQ ID NO:16:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: $6..1\overline{1}$
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 12..26
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:16:

CGACACTCGA GGTGACGGAC AAGGTC

- (2) INFORMATION FOR SEQ ID NO:17:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 12..26
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:17:

CGACAGTCGA CCCAATCAGG GACCTC

26

- (2) INFORMATION FOR SEQ ID NO:18:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 10..15
 - (D) OTHER INFORMATION: /note= "Bsi WI restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 6..32
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:18:

TCGAG TAT CCG TAC GAC GTA CCA GAC TAC GCA G

Tyr Pro Tyr Asp Val Pro Asp Tyr Ala

1

- (2) INFORMATION FOR SEQ ID NO:19:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 9 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:19:

Tyr Pro Tyr Asp Val Pro Asp Tyr Ala

- (2) INFORMATION FOR SEQ ID NO:20:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:20:

TCGACTGCGT AGTCTGGTAC GTCGTACGGA TAC

33

- (2) INFORMATION FOR SEQ ID NO:21:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:21:

TCGACTATCC GTACGACGTA CCAGACTACG CAC

- (2) INFORMATION FOR SEQ ID NO:22:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 33 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:22:

TCGAGTGCGT AGTCTGGTAC GTCGTACGGA TAG

- (2) INFORMATION FOR SEQ ID NO:23:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 80 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Sac II restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: 12..16
 - (D) OTHER INFORMATION: /note= "Kozak sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: 17..58
 - (D) OTHER INFORMATION: /note= "Myristoylation signal."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 59..64
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 65..80
 - (D) OTHER INFORMATION: /note= "Zeta homology."
 - (ix) FEATURE:

- (A) NAME/KEY: CDS (B) LOCATION: 17..79
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:23:

CGACACCGCG GCCACC ATG GGG AGT AGC AAG AGC AAG CCT AAG GAC CCC 49 Met Gly Ser Ser Lys Ser Lys Pro Lys Asp Pro

AGC CAG CGC CTC GAG AGG AGT GCA GAG ACT G Ser Gln Arg Leu Glu Arg Ser Ala Glu Thr

80

- (2) INFORMATION FOR SEQ ID NO: 24:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 21 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:24:

Met Gly Ser Ser Lys Ser Lys Pro Lys Asp Pro Ser Gln Arg Leu Glu 10

Arg Ser Ala Glu Thr 20

- (2) INFORMATION FOR SEQ ID NO:25:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 12..26
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: $12..\overline{27}$

- (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:25:

CGACACTCGA G GAG CTC TGT GAC GAT G Glu Leu Cys Asp Asp

27

- (2) INFORMATION FOR SEQ ID NO:26:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:26:

Glu Leu Cys Asp Asp 1

- (2) INFORMATION FOR SEQ ID NO:27:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 41 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single

 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 12..41
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 27..29
 - (D) OTHER INFORMATION: /note= "GAT to AAG."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 9..41

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:27:

CGACACTC GAG CTC TGC TAC TTG CTA AAG GGA ATC CTC TTC Glu Leu Cys Tyr Leu Leu Lys Gly Ile Leu Phe 1 5 10

41

- (2) INFORMATION FOR SEQ ID NO:28:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:28:

Glu Leu Cys Tyr Leu Leu Lys Gly Ile Leu Phe 1 5 10

- (2) INFORMATION FOR SEQ ID NO:29:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 44 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single'
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $6..1\overline{1}$
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 9.44
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 27..44
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:29:

CGACACTC GAG CTG CTG GAT CCG AAG CTC TGC TAC TTG CTA AAG
Glu Leu Leu Asp Pro Lys Leu Cys Tyr Leu Leu Lys
1 5 10

- (2) INFORMATION FOR SEQ ID NO:30:
 - (i) SEQUENCE CHARACTERISTICS:

31

- (A) LENGTH: 12 amino acids
- (B) TYPE: amino acid
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: protein
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:30:

Glu Leu Leu Asp Pro Lys Leu Cys Tyr Leu Leu Lys

- (2) INFORMATION FOR SEQ ID NO:31:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 31 base pairs

 - (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $6..1\overline{1}$
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $12..\overline{31}$
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 9..31
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:31:

CGACACTC GAG ACA ACA GAG TAC CAG GTA GC Glu Thr Thr Glu Tyr Gln Val Ala 1

- (2) INFORMATION FOR SEQ ID NO:32:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 8 amino acids
 - (B) TYPE: amino acid (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:32:

Glu Thr Thr Glu Tyr Gln Val Ala

- (2) INFORMATION FOR SEQ ID NO:33:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: $12..\overline{28}$
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 9..28
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:33:

CGACACTC GAG GGC GTG CAG GTG GAG AC
Glu Gly Val Gln Val Glu Thr

28

- (2) INFORMATION FOR SEQ ID NO:34:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:34:

Glu Gly Val Gln Val Glu Thr

- (2) INFORMATION FOR SEQ ID NO:35:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single

27

- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $6..1\overline{1}$
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 12..27
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: complement (9..26)
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:35:

CGACAGTCGA CTTCCAGTTT TAGAAGC

- (2) INFORMATION FOR SEQ ID NO:36:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:36:

Leu Leu Lys Leu Glu Val 1

- (2) INFORMATION FOR SEQ ID NO:37:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid

 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 7..12
 - (D) OTHER INFORMATION: /note= "Xho I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS

(B) LOCATION: 10..27

(ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: 13..27
- (D) OTHER INFORMATION: /note= "Region of homology with target sequence."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:37:

TCGACACTC GAG ACG GGG GCC GAG GGC
Glu Thr Gly Ala Glu Gly
1 5

27

- (2) INFORMATION FOR SEQ ID NO:38:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:38:

Glu Thr Gly Ala Glu Gly

- (2) INFORMATION FOR SEQ ID NO:39:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 7..12
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: complement (10..18)
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: $13..\overline{2}8$
 - (D) OTHER INFORMATION: /note= "Region of homology with target sequence."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:39:	
CCGACAGTCG ACCTCTATTT TGAGCAGC	28
(2) INFORMATION FOR SEQ ID NO:40:	
(i) SEQUENCE CHARACTERISTICS:(A) LENGTH: 3 amino acids(B) TYPE: amino acid(D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: protein	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:40:	
Ile Glu Val 1	
(2) INFORMATION FOR SEQ ID NO:41:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 38 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: CDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:41:	
CGACACCGCG GCCACCATGA AGCTACTGTC TTCTATCG	38
(2) INFORMATION FOR SEQ ID NO:42:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:42:	
CGACAGTCGA CCGATACAGT CAACTGTC	28
(2) INFORMATION FOR SEQ ID NO:43:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 38 base pairs	

- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Sac II restriction site."
- (ix) FEATURE:
 - (A) NAME/KEY: misc signal
 - (B) LOCATION: $12..\overline{16}$
 - (D) OTHER INFORMATION: /note= "Kozak sequence."
- (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 17..37
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 17..38
 - (D) OTHER INFORMATION: /note= "Gal4 (1-147) coding region."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:43:

CGACACCGCG GCCACC ATG AAG CTA CTG TCT TCT ATC G Met Lys Leu Leu Ser Ser Ile

38

- (2) INFORMATION FOR SEQ ID NO:44:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 7 amino acids

 - (B) TYPE: amino acid (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:44:

Met Lys Leu Leu Ser Ser Ile 5

- (2) INFORMATION FOR SEQ ID NO:45:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:

- (A) NAME/KEY: misc_feature
- (B) LOCATION: 1..17
- (D) OTHER INFORMATION: /note= "Region encoding for C-terminal end of Gal4 (1-147)."
- (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 3..17
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 18..23
 - (D) OTHER INFORMATION: /note= "Sal I restriction site."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:45:
- GA CAG TTG ACT GTA TCG GTCGACTGTC G
 Arg Gln Leu Thr Val Ser
 1 5

28

- (2) INFORMATION FOR SEQ ID NO:46:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:46:

Arg Gln Leu Thr Val Ser

- (2) INFORMATION FOR SEQ ID NO:47:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 34 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:47:

CGACACCGCG GCCACCATGG TTTCTAAGCT GAGC

- (2) INFORMATION FOR SEQ ID NO:48:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs

- (B) TYPE: nucleic acid
- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:48:

CGACAGTCGA CCAACTTGTG CCGGAAGG

28

- (2) INFORMATION FOR SEQ ID NO:49:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 34 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 6..11
 - (D) OTHER INFORMATION: /note= "Sac II restriction site."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: 12..16
 - (D) OTHER INFORMATION: /note= "Kozak sequence."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 17..34
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 17..34
 - (D) OTHER INFORMATION: /note= "Region encoding N-terminal end of HNF1 (1281)."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:49:

CGACACCGCG GCCACC ATG GTT TCT AAG CTG AGC
Met Val Ser Lys Leu Ser
1

- (2) INFORMATION FOR SEQ ID NO:50:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein

- (ii) MOLECULE TYPE: cDNA
- (ix) FEATURE:
 - (A) NAME/KEY: misc_signal
 - (B) LOCATION: 3..7
 - (D) OTHER INFORMATION: /note= "Kozak sequence."
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 1..11
 - (D) OTHER INFORMATION: /note= "Complementary to bases 5 to 15 of SEQ ID NO:54."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:53:

GGCCACCATG C

11

- (2) INFORMATION FOR SEQ ID NO:54:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 3 amino acids

 - (B) TYPE: amino acid
 (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..3
 - (D) OTHER INFORMATION: /note= "Translation product of SEQ ID NO:53 and SEQ ID NO:55. Translational start site at base 8 of SEQ ID NO:53."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:54:

Met Leu Glu 1

- (2) INFORMATION FOR SEQ ID NO:55:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 17 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 14..17
 - (D) OTHER INFORMATION: /note= "Sac II restriction site overhang."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:50:

Met Val Ser Lys Leu Ser 1 5

- (2) INFORMATION FOR SEQ ID NO:51:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..20
 - (D) OTHER INFORMATION: /note= "Region encoding for C-terminal end of HNF1 (1-282)."
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 3..17
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:51:

CC TTC CGG CAC AAG TTG GTCGACTGTC G Ala Phe Arg His Lys Leu 1 5 28

- (2) INFORMATION FOR SEQ ID NO:52:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:52:

Ala Phe Arg His Lys Leu
1 5

- (2) INFORMATION FOR SEQ ID NO:53:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear

- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Sal I restriction site overhang."
- (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 5..27
 - (D) OTHER INFORMATION: /note= "Complementary to SEQ ID NO:60, bases 5 to 27."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:58:

TCGACCCTAA GAAGAAGAGA AAGGTAC

27

- (2) INFORMATION FOR SEQ ID NO:59:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ix) FEATURE:
 - (A) NAME/KEY: Peptide
 - (B) LOCATION: 1..11
 - (D) OTHER INFORMATION: /note= "Translation product of SEQ ID NOS:58 and 60."
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:59:

Leu Asp Pro Lys Lys Lys Arg Lys Val Leu Glu
1 5 10

- (2) INFORMATION FOR SEQ ID NO:60:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 27 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: both
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: misc feature
 - (B) LOCATION: 1..5
 - (D) OTHER INFORMATION: /note= "Xho I restriction site overhang."
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature

	(ix)	FEATURE:	
		(A) NAME/KEY: misc_feature (B) LOCATION: 15	
		(D) OTHER INFORMATION: /note= "Xho I restriction site overha	ing."
	(ix)	FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 515 (D) OTHER INFORMATION: /note= "Complementary to bases 1 to 1 SEQ ID NO:53."	.1 of
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:55:	
TCG	GCAT	GG TGGCCGC	17
(2)	INFO	RMATION FOR SEQ ID NO:56:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: CDNA	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:56:	
TCG	ACCCT	'AA GAMGAAGAGA AAGGTAC	27
(2)	INFO	RMATION FOR SEQ ID NO:57:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: CDNA	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:57:	
TCG	AGTAC	CT TTCTCTTCKT CTTAGGG	27
(2)	INFO	RMATION FOR SEQ ID NO:58:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: both (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: cDNA	

		(B) LOCATION: 527 (D) OTHER INFORMATION: /note= "Complementary to SEQ ID NO:58, bases 5 to 27."	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:60:	
TCGA	AGTAC	CT TTCTCTTCTT CTTAGGG	27
(2)	INFO	RMATION FOR SEQ ID NO:61:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: cDNA	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:61:	
CGA	CAGTC	GA CGCCCCCCG ACCGATGTC	29
(2)	INFO	RMATION FOR SEQ ID NO:62:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 26 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: cDNA	
	(xi)	SEQUENCE DESCRIPTION: SEQ ID NO:62:	
CGA	CACTC	GA GCCCACCGTA CTCGTC	26
(2)	INFO	RMATION FOR SEQ ID NO:63:	
	(i)	SEQUENCE CHARACTERISTICS: (A) LENGTH: 29 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
	(ii)	MOLECULE TYPE: cDNA	
	(ix)	<pre>FEATURE: (A) NAME/KEY: misc_feature (B) LOCATION: 611 (D) OTHER INFORMATION: /note= "Sal I restriction site."</pre>	
	(ix)	FEATURE:	

- (A) NAME/KEY: CDS
- (B) LOCATION: 12..29
- (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 12..29
 - (D) OTHER INFORMATION: /note= "Region encoding N^^terminal end of VP16 (413^^490)."
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:63:

CGACAGTCGA C GCC CCC CCG ACC GAT GTC
Ala Pro Pro Thr Asp Val
1

29

- (2) INFORMATION FOR SEQ ID NO:64:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 6 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:64:

Ala Pro Pro Thr Asp Val

- (2) INFORMATION FOR SEQ ID NO:65:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: double
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (ix) FEATURE:
 - (A) NAME/KEY: CDS
 - (B) LOCATION: 1..15
 - (ix) FEATURE:
 - (A) NAME/KEY: misc_feature
 - (B) LOCATION: 1..15
 - (D) OTHER INFORMATION: /note= "Region encoding C-terminal end of VP16 (413-490)."

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:65:

GAC GAG TAC GGT GGG CTCGAGTGTC G Asp Glu Tyr Gly Gly 1

26

- (2) INFORMATION FOR SEQ ID NO:66:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5 amino acids
 (B) TYPE: amino acid
 (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: protein
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:66:

Asp Glu Tyr Gly Gly

- (2) INFORMATION FOR SEQ ID NO:67:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 23 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:67:

GGAATTCCAT ATGGGCGTGC AGG

23

- (2) INFORMATION FOR SEQ ID NO:68:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 5 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:68:

His Met Gly Val Gln 1

- (2) INFORMATION FOR SEQ ID NO:69:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 39 base pairs
 - (B) TYPE: nucleic acid

- (C) STRANDEDNESS: single
- (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:69:

CTGTCCCGGG ANNNNNNNN TTTCTTTCCA TCTTCAAGC

39

- (2) INFORMATION FOR SEQ ID NO:70:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 11 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:70:

Arg Ser Xaa Xaa Xaa Lys Lys Gly Asp Glu Leu 1 5 10

- (2) INFORMATION FOR SEQ ID NO:71:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 64 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:71:

CTGTCCCGGG AGGAATCAAA TTTCTTTCCA TCTTCAAGCA TNNNNNNNNN GTGCACCACG 60

- (2) INFORMATION FOR SEQ ID NO:72:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 19 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEQ ID NO:72:

Arg Ser Ser Asp Phe Lys Lys Gly Asp Glu Leu Met Xaa Xaa Xaa His 1 5 10 15

Val Val Cys

57

28

- (2) INFORMATION FOR SEQ ID NO:73:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 57 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:73:

CGCGGATCCT CATTCCAGTT TTAGAAGCTC CACATCNNNN NNNNNAGTGG CATGTGG

- (2) INFORMATION FOR SEQ ID NO:74:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 15 amino acids
 - (B) TYPE: amino acid
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide
 - (xi) SEQUENCE DESCRIPTION: SEO ID NO:74:
- Glu Leu Lys Leu Leu Glu Val Asp Xaa Xaa Xaa Thr Ala His Pro
- (2) INFORMATION FOR SEQ ID NO:75:
 - (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 28 base pairs
 - (B) TYPE: nucleic acid
 - (C) STRANDEDNESS: single
 - (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: CDNA
 - SEQUENCE DESCRIPTION: SEQ ID NO:75:

CGCGGATCCT CATTCCAGTT TTAGAAGC

(2) INFORMATION FOR SEQ ID NO:76:

- - (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 5 amino acids
 - (B) TYPE: amino acid (D) TOPOLOGY: linear
 - (ii) MOLECULE TYPE: peptide

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:76:	
Glu Leu Lys Leu 1 5	
(2) INFORMATION FOR SEQ ID NO:77:	
 (i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear 	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:77:	
CGACAGTCGA CCGATACAGT CAACTGTC	2.8
(2) INFORMATION FOR SEQ ID NO:78:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 28 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:78:	
CGACAGTCGA CCAACTTGTG CCGGAAGG	28
(2) INFORMATION FOR SEQ ID NO:79:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 17 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single (D) TOPOLOGY: linear	
(ii) MOLECULE TYPE: cDNA	
(xi) SEQUENCE DESCRIPTION: SEQ ID NO:79:	
TCGAGCATGG TGGCCGC	17
(2) INFORMATION FOR SEQ ID NO:80:	
(i) SEQUENCE CHARACTERISTICS: (A) LENGTH: 27 base pairs (B) TYPE: nucleic acid (C) STRANDEDNESS: single	

(D) TOPOLOGY:	linear
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- (ii) MOLECULE TYPE: cDNA
- SEQUENCE DESCRIPTION: SEQ ID NO:80:

TCGAGTACCT TTCTCTTCTT CTTAGGG

27

(2) INFORMATION FOR SEQ ID NO:81:

- (i) SEQUENCE CHARACTERISTICS:
 - (A) LENGTH: 26 base pairs
 (B) TYPE: nucleic acid
 (C) STRANDEDNESS: single
 (D) TOPOLOGY: linear
- (ii) MOLECULE TYPE: cDNA
- (xi) SEQUENCE DESCRIPTION: SEQ ID NO:81:

CGACACTCGA GCCCACCGTA CTCGTC